

APPROVED	O.G. FIG.	
DRAFTSMAN	CLASS	SUBCLASS

6007988

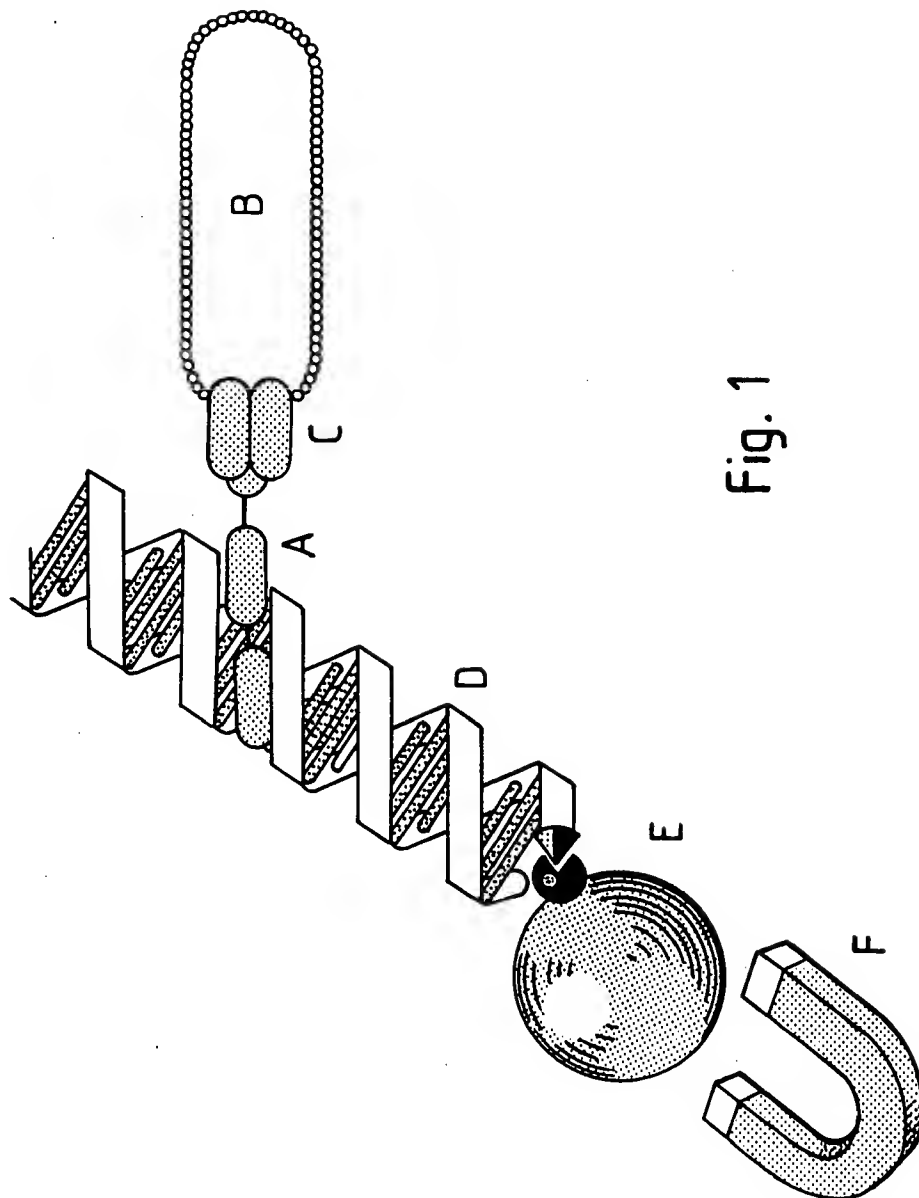


Fig. 1

APPROVED	O.G. FIG.	
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-1123456789

MAERP⊙A⊙PVES⊙DRR⊙SRSD⊙ELTR⊙TRI⊙T
 GQK⊙P⊙QQ⊙RI--⊙MRN⊙SXX⊙XX⊙X^R⊙X^K⊙T⊙T
 GEK⊙P⊙A⊙DI--⊙GRK⊙ARSDERK⊙R⊙T⊙KI⊙HLRQKD

$$\frac{\beta}{\beta} \quad \frac{\alpha}{\alpha}$$

Fig. 2

APPROVED	O.G. FIG.	
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(i) TATGACTTGGATGGGAGACCGCCTGG
 ACTGAACCTACCCCTCTGGCGGACCTTAA - (B)

(ii) TATATAGCGTGGCGGTATATA
 ATATATCGCACCGGCATATATGCG - (B)

(iii) TATATAGCGGXXGCGGTATATA
 ATATATCGGXXGCGCATATATGCG - (B)

Fig. 3

APPROVED	O.G. FIG.	
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OLIGONUCLEOTIDE LIBRARY

Fig. 4

		5'	G	A	T	C	N	N	N	N	N	N	N	N	N	3'
		↓	N	N	N	N	G	A	T	C	N	N	N	N	N	
		3'	N	N	N	N	N	N	N	N	G	A	T	C	5'→3'	
α-HELIX SEQUENCE	123456789	•RSDHLTTHIR														G _T TGG
		RVDALAHRR														GTG
		QRASLASHMR														G _T TA
		NRDTLTRHSK														GTT
		QKGHLTEHRK														GGA
		QSVHLQSHSR														TGA
		RLDGLRTHLK														G _T A _{T/C} G
		TPGNLTRHGR														GAT
		NGGNLGRHMK														GAT
		RADALMVHKR														G _T TG
		NQSNLERHHR														GAT
		DRSNLERHTR														GAC
		RSDTLKKGK														GCG
		QQSNLVRHQR														GAT
		NGANLERHRR														GAT
		RGDALTSHER														G _T TG
		RGDHLKDHIK														G _T GG
		RGPDLARHGR														GCG
		REDVLI RHGK														GT _C G
		RSDDLQRHHK														GT _C G
		RQDTLVGHER														G _T _{T/C} G
		RAADLNHVR														GCG
		SQGNLQRHGR														GA _{G/T}
		ITGGSLARHER														GTT
		DHANLARHTR														GAC
		LQSNLVRHQR														GAT _C
		RKDVLVSHVR														G _T _{T/C} TG
		RRDVLMMHIR														GT _C G
		QGGNLVRHLR														GAA
		SRDVLRRHNR														GCT
		EKATLARHMK														GCT
		QAQTLQRHLK														GCT
		IASNLLRHQR														GA _T _{A/C}

BINDING SITE SIGNATURE

APPROVED	O.G. FIG.	
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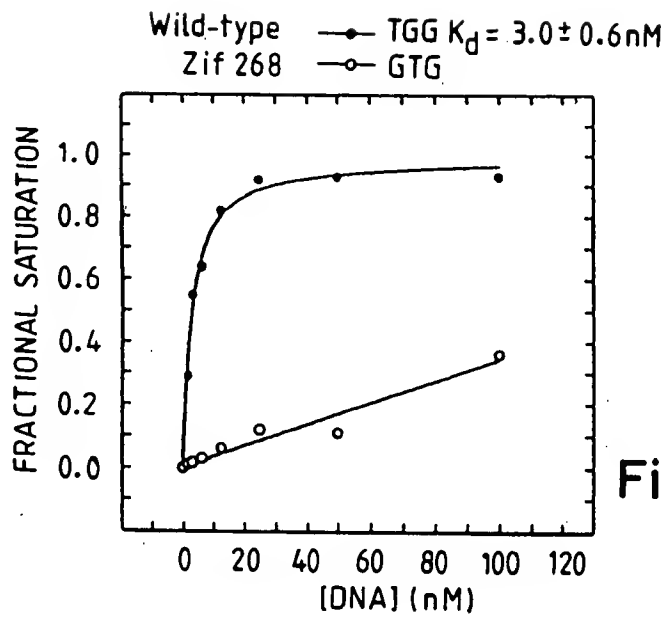


Fig. 5A

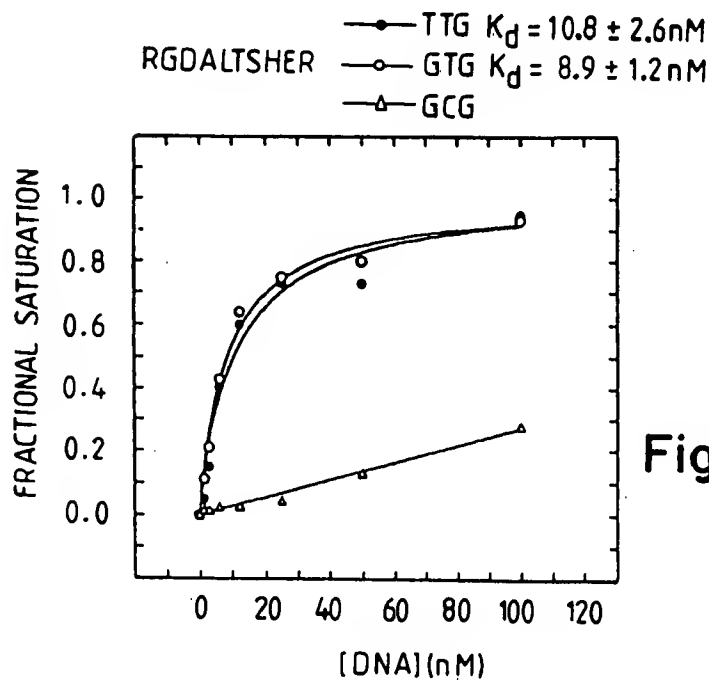


Fig. 5B

APPROVED	O.G. FIG.	
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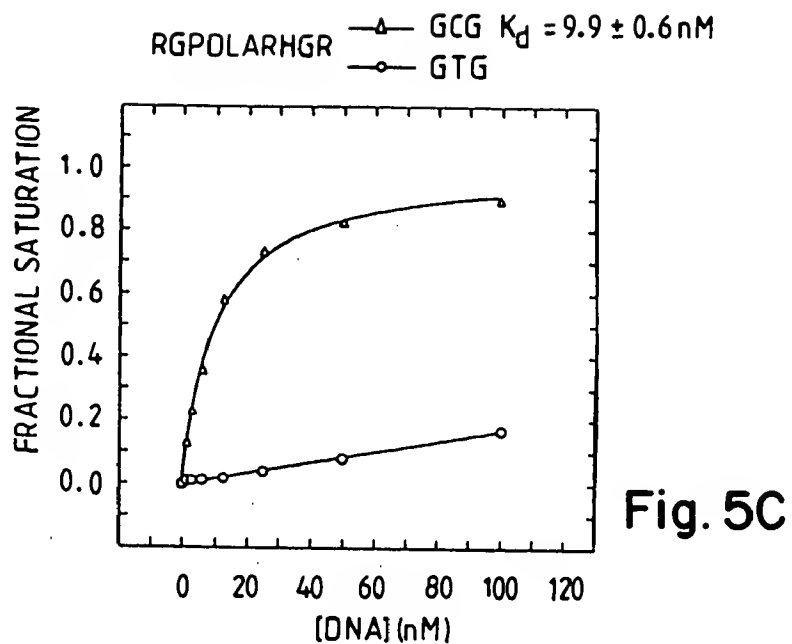


Fig. 5C

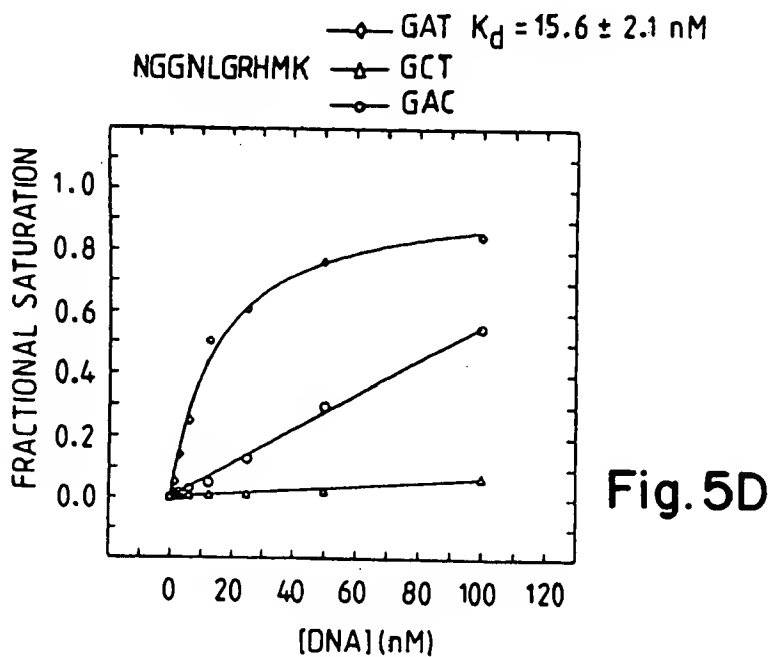


Fig. 5D

APPROVED	O.G. FIG.	
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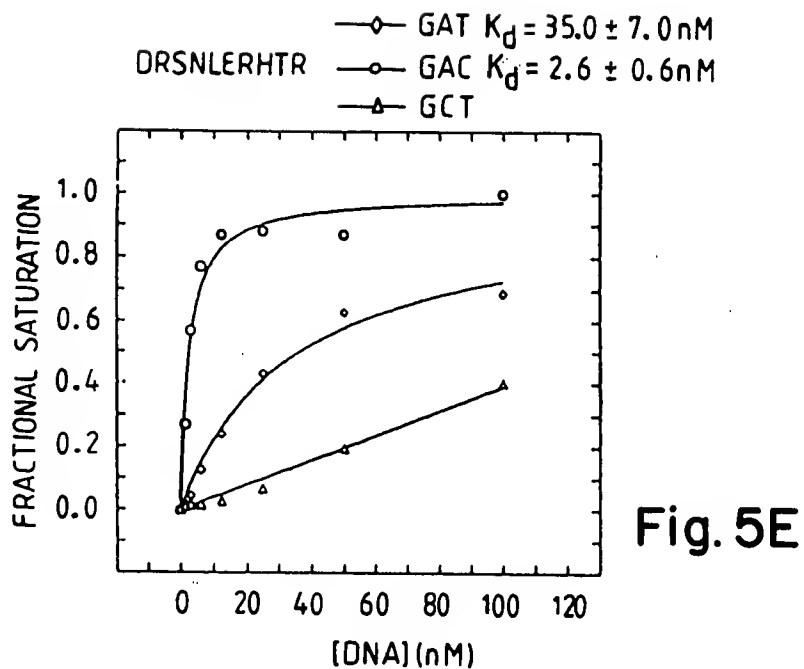


Fig. 5E

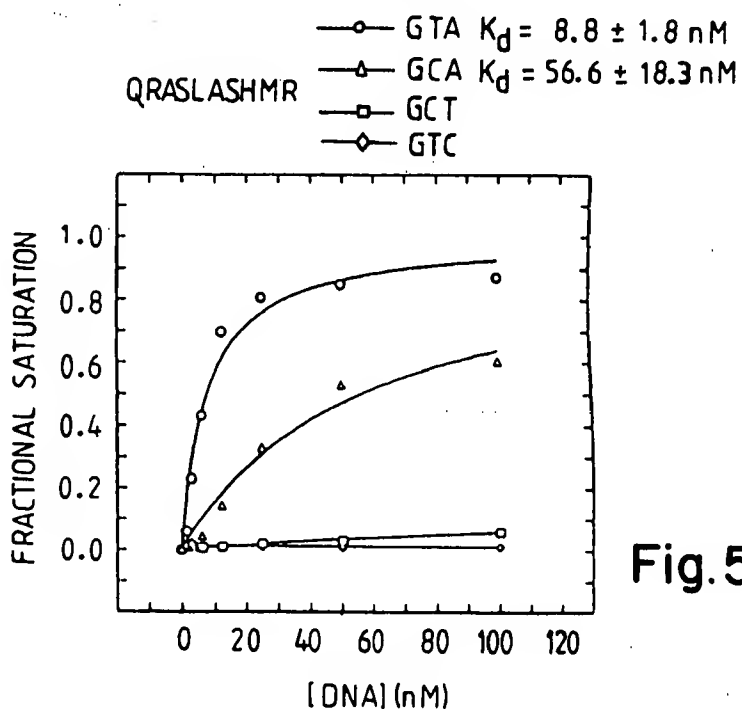


Fig. 5F

APPROVED	O.G. FIG.	
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1 *BCR - ABL* TTC CAT GGA GAC GCA G AA GCC CTT CAG CGG CCA

2 *BCR* TTC CAT GGA GAC GCA G gt gag ttc ctc acg cca

3 *ABL* ccc ctt tct ctt cca g AA GCC CTT CAG CGG CCA

Fig.6

APPROVED	O.G. FIG.	
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		-1	1	2	3	4	5	6	7	8	9	
1A	MAEEKPFQ	CRI	CMRN	FSD	RSSL	TR	H	TR	H			TGEKP
1B	MAEEKPFQ	CRI	CMRN	FSE	RGT	LA	R	H	E	K	H	TGEKP
2A		FQ	CRI	CMRN	FSG	GNL	VR	H	L	R	H	TGEKP
3A		FQ	CRI	CMRN	FSA	QTL	QR	H	L	K	H	TGEK
3B		FQ	CRI	CMRN	FSA	ATL	QR	H	L	K	H	TGEK
3C		FQ	CRI	CMRN	FSA	QDL	QR	H	L	K	H	TGEK
	β -SHEET		β -SHEET		α -HELIX							LINKER

Fig. 7

APPROVED	O.G. FIG.	
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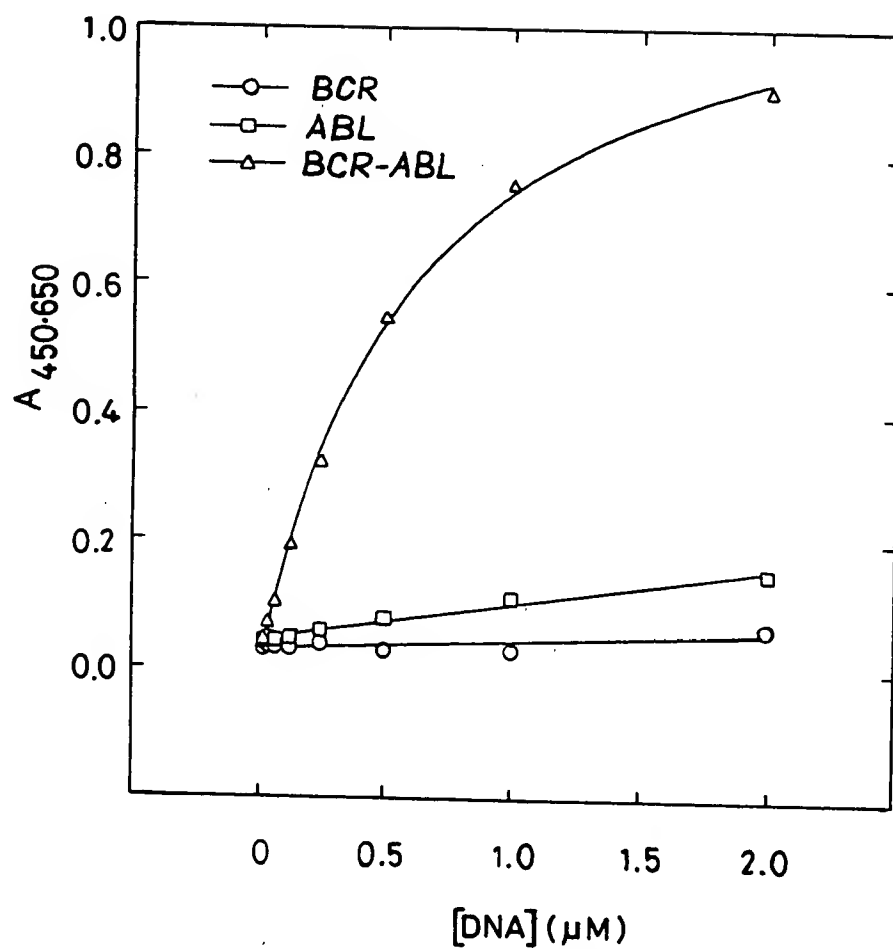


Fig. 8

APPROVED	O.G. FIG.	
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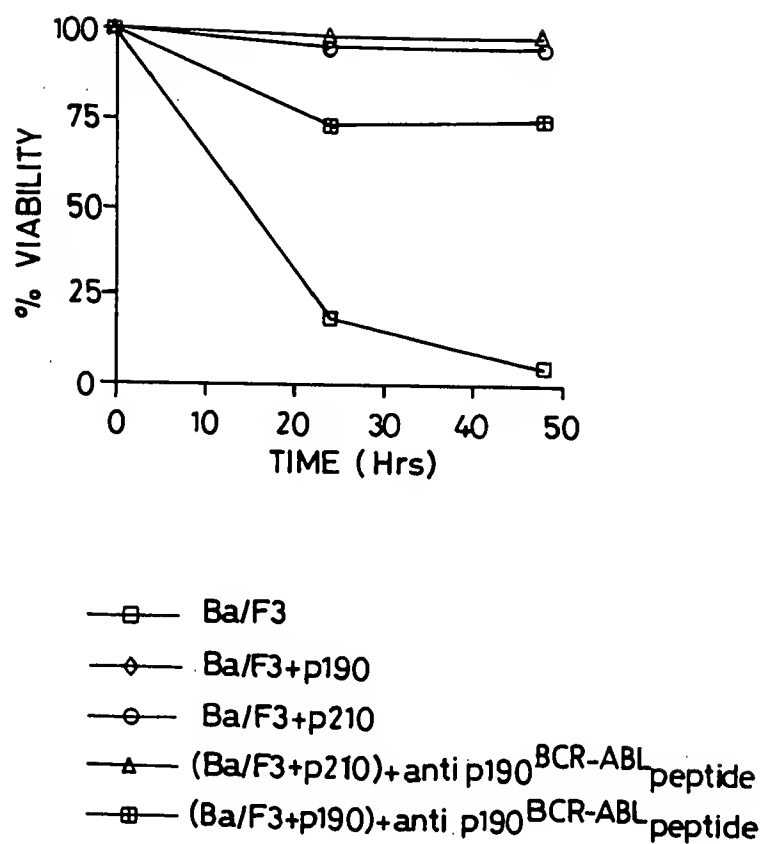


Fig. 9

APPROVED	O.G. FIG.	
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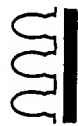


Fig. 10A

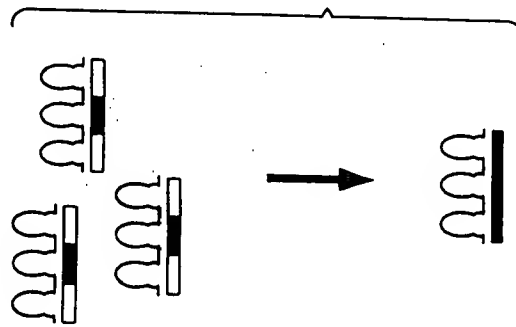


Fig. 10B

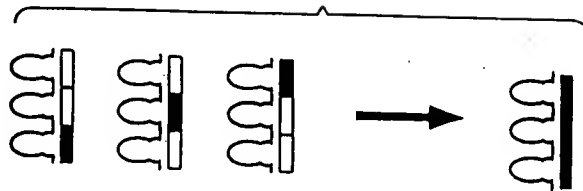


Fig. 10C

APPROVED	O.G. FIG.	
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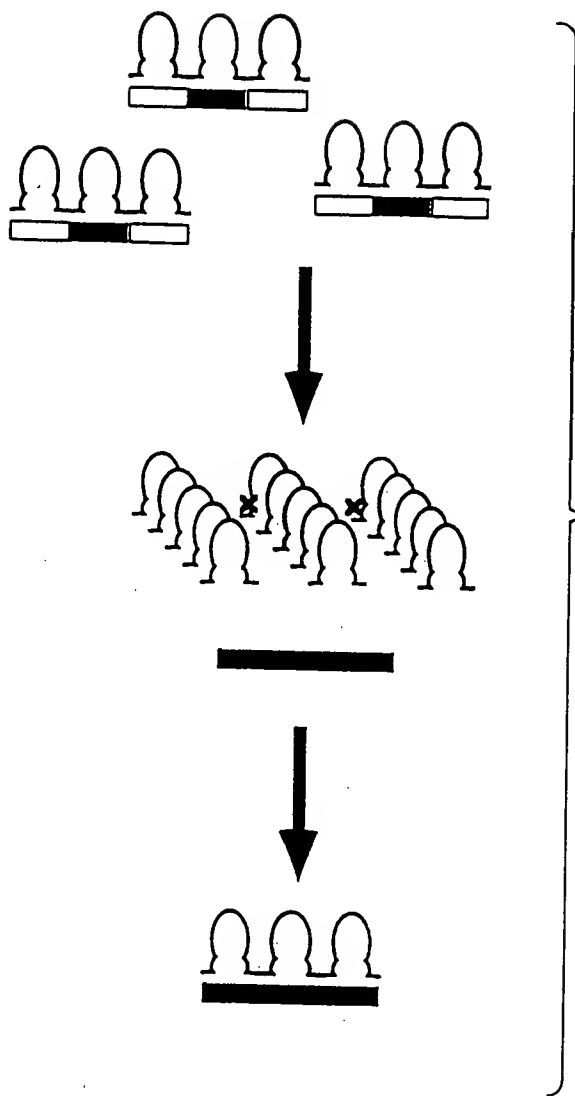


Fig. II

APPROVED	O.G. FIG.	
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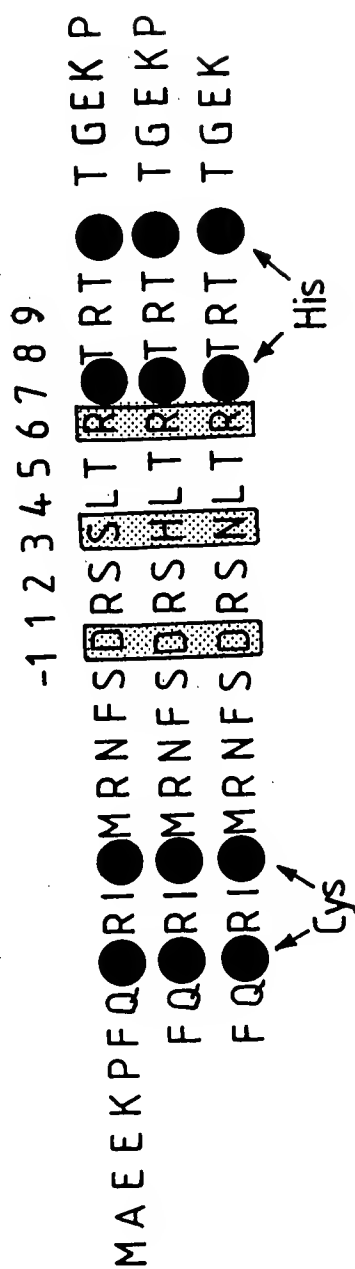


Fig. 12

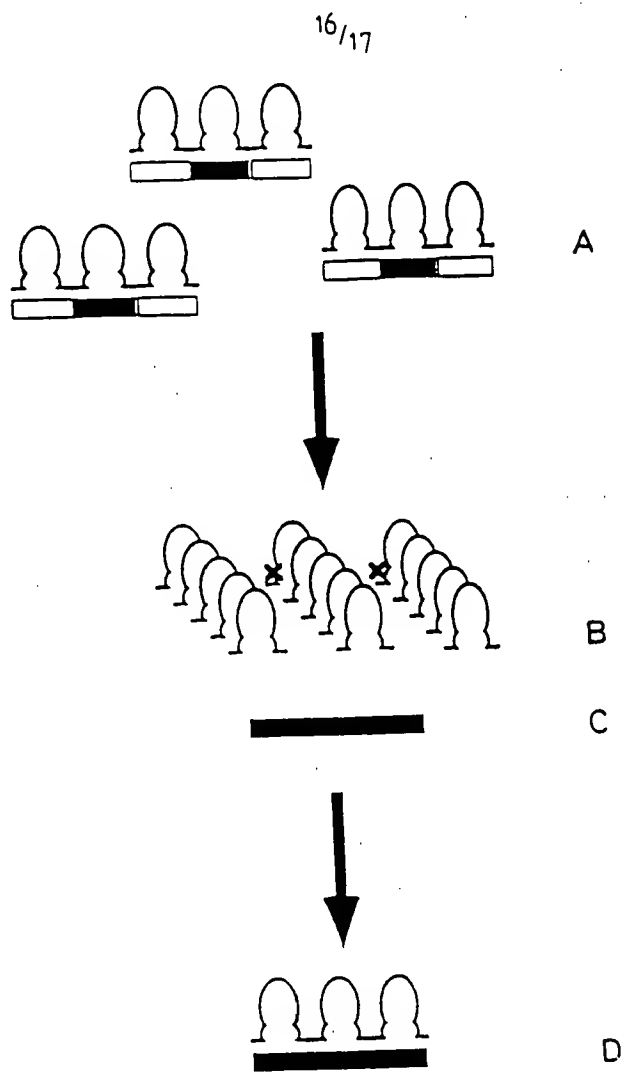


Fig. 14